

<110> DeAngelis, Paul Jing, Wei

<120> TARGETED GLYCOSAMINOGLYCAN POLYMERS BY POLYMER GRAFTING AND METHODS OF MAKING AND USING SAME

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<213> Pasteurella multocida

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Val Asn Ala Leu Leu Ser Asn Lys Asp Asn Ile Ala Val Arg Cys Ala

Page 23

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<213> Pasteurella multocida

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1320

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1800

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Page 46

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Page 50

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1860

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1860

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1860

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<211> 2007

<212> DNA

<213> Pasteurella multocida

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1680

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Walter State Control

<211> 501

<212> PRT

<213> Pasteurella multocida

<400> 61

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Gln Ser Lys Ser Asn Lys Ile Glu Glu Asp Asn Ile Ser Gly Glu Asn 50 55 60

Glu Phe Ser Val Ser Ile Lys Asp Leu Tyr Asn Glu Ile Ser Asn Ser 65 70 75 80

Glu Leu Gly Ile Thr Lys Glu Arg Leu Gly Ala Pro Pro Leu Val Ser 85 90 95

Ile Ile Met Thr Ser His Asn Thr Glu Lys Phe Ile Glu Ala Ser Ile 100 105 110

Asn Ser Leu Leu Gln Thr Tyr Asn Asn Leu Glu Val Ile Val Val 115 120 125

Asp Asp Tyr Ser Thr Asp Lys Thr Phe Gln Ile Ala Ser Arg Ile Ala 130 135 140

Asn Ser Thr Ser Lys Val Lys Thr Phe Arg Leu Asn Ser Asn Leu Gly 145 150 155 160

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Thr	Tyr	Phe	Ala	Lys 165	Asn	Thr	Gly	Ile	Leu 170	Lys	Ser	Lys	Gly	Asp 175	Ile
Ile	Phe	Phe	Gln 180	Asp	Ser	Asp	Asp	Val 185	Cys	His	His	Glu	Arg 190	Ile	Glu
Arg	Cys	Val 195	Asn	Ala	Leu	Leu	Ser 200	Asn	Lys	Asp	Asn	Ile 205	Ala	Val	Arg
Cys	Ala 210	Tyr	Ser	Arg	Ile	Asn 215	Leu	Glu	Thr	Gln	Asn 220	Ile	Ile	Lys	Val
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Lys	Val	Phe	Asn	Glu 245	Ile	Gly	Phe	Phe	Asn 250	Cys	Thr	Thr	Lys	Ala 255	Ser
Asp	Asp	Glu	Phe 260	Tyr	His	Arg	Ile	Ile 265	Lys	Tyr	Tyr	Gly	Lys 270	Asn	Arg
Ile	Asn	Asn 275	Leu	Phe	Leu	Pro	Leu 280	Tyr	Tyr	Asn	Thr	Met 285	Arg	Glu	Asp
Ser	Leu 290	Phe	Ser	Asp	Met	Val 295	Glu	Trp	Val	Asp	Glu 300	Asn	Asn	Ile	Lys
Gln 305	Lys	Thr	Ser	Asp	Ala 310	Arg	Gln	Asn	Tyr	Leu 315	His	Glu	Phe	Gln	Lys 320
Ile	His	Asn	Glu	Arg 325	Lys	Phe	Asn	Glu	Leu 330	Lys	Glu	Ile	Phe	Ser 335	Phe
Pro	Arg	Ile	His 340	Asp	Ala	Leu		Ile 345 Page		Lys	Glu	Met	Ser 350	Lys	Leu

Ser Asn Pro Lys Ile Pro Val Tyr Ile Asn Ile Cys Ser Ile Pro Ser 355 360 365

Arg Ile Lys Gln Leu Gln Tyr Thr Ile Gly Val Leu Lys Asn Gln Cys 370 375 380

Asp His Phe His Ile Tyr Leu Asp Gly Tyr Pro Glu Val Pro Asp Phe 385 390 395 400

Ile Lys Lys Leu Gly Asn Lys Ala Thr Val Ile Asn Cys Gln Asn Lys
405 410 415

Asn Glu Ser Ile Arg Asp Asn Gly Lys Phe Ile Leu Leu Glu Lys Leu 420 425 430

Ile Lys Glu Asn Lys Asp Gly Tyr Tyr Ile Thr Cys Asp Asp Asp Ile 435 440 445

Arg Tyr Pro Ala Asp Tyr Ile Asn Thr Met Ile Lys Lys Ile Asn Lys 450 455 460

Tyr Asn Asp Lys Ala Ala Ile Gly Leu His Gly Val Ile Phe Pro Ser 465 470 475 480

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ttctggagaa 240	aacgaatttt	cagtatcaat	aaaagatcta	tataacgaaa	taagcaatag
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 105 110

Arg Glu Val Ser Ser Gly Ser Met Glu Gly Leu Trp Asn Thr Phe Thr 195 200 205

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Tyr Ser Lys Leu Asn Leu Glu Leu Val Tyr Asn Val Glu Gly 225 230 235

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<212> PRT

<213> Escherichia coli

<400> 64

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Val Asp Leu Gln His Lys Asn Ser Pro Leu Lys Gly Asn Asp Asn Leu 35 40 45

Ile His Lys Arg Ile Asn Glu Tyr Asp Asn Val Leu Glu Leu Ser Lys 50 55 60

Asn Val Ser Ala Gln Asn Ser Gly Asn Glu Phe Ser Tyr Leu Leu Gly 65 70 75 80

Tyr Ala Asp Ser Leu Arg Lys Val Gly Met Leu Asp Thr Tyr Ile Lys 85 90 95

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Arg	Val	Lys	Leu	Phe	Glu	His	Ile	Ser	Asn	Ala	Leu	Arg	Tyr	Ser	Arg
		115					120					125			

Asn	Ala	Asn	Asp	Asp	Met	Gln	Asp	Ser	Tyr	Asn	Leu	Leu	Pro	Glu	Gln
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Glu Le	ı Phe	Ile	Cys	Asn	Ala	Val	Ile	Asp	Asn	Met	Asn	Ile	Tyr	Arg
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Ser	Gln	Phe	Asn	Lys	Cys	Leu	Arg	Lys	Tyr	Asp	Leu	Ser	Glu	Ile	Thr
225					230					235					240

Phe Asn Ser Glu Asp Thr Ile Ala Tyr Ser Leu His Ser Leu Leu Asn 275 280 285

SEQUENCE LISTING.ST25.txt Gln Thr Tyr Glu Asn Ile Glu Ile Leu Val Cys Asp Asp Cys Ser Ser 290 295 300 Asp Lys Ser Leu Glu Ile Ile Lys Ser Ile Ala Tyr Ser Ser Ser Arg

310

305

Val Lys Val Tyr Ser Ser Arg Lys Asn Gln Gly Pro Tyr Asn Ile Arg 325 330 335

315

320

Asn Glu Leu Ile Lys Lys Ala His Gly Asn Phe Ile Thr Phe Gln Asp 340 345 350

Ala Asp Asp Leu Ser His Pro Glu Arg Ile Gln Arg Gln Val Glu Val 355 360 365

Leu Arg Asn Asn Lys Ala Val Ile Cys Met Ala Asn Trp Ile Arg Val 370 375 380

Ala Ser Asn Gly Lys Ile Gln Phe Phe Tyr Asp Asp Lys Ala Thr Arg 385 390 395 400

Met Ser Val Val Ser Ser Met Ile Lys Lys Asp Ile Phe Ala Thr Val 405 410 415

Gly Gly Tyr Arg Gln Ser Leu Ile Gly Ala Asp Thr Glu Phe Tyr Glu 420 425 430

Thr Val Ile Met Arg Tyr Gly Arg Glu Ser Ile Val Arg Leu Leu Gln 435 440 445

Pro Leu Ile Leu Gly Leu Trp Gly Asp Ser Gly Leu Thr Arg Asn Lys 450 455 460

Gly Thr Glu Ala Leu Pro Asp Gly Tyr Ile Ser Gln Ser Arg Arg Glu 465 470 475 480

Tyr Ser Asp Ile Ala Ala Arg Gln Arg Val Leu Gly Lys Ser Ile Val 485 490 495

Ser Asp Lys Asp Val Arg Gly Leu Leu Ser Arg Tyr Gly Leu Phe Lys 500 505 510

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<212> PRT

<213> Mus musculus

<400> 65

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Ser His Ser Arg Arg Glu Glu His Ser Gly Arg Asn Gly Leu His Gln
35 40 45

Pro Ser Pro Asp His Phe Trp Pro Arg Phe Pro Asp Ala Leu Arg Pro 50 55 60

Phe Phe Pro Trp Asp Gln Leu Glu Asn Glu Asp Ser Ser Val His Ile 70 75 80

Ser Pro Arg Gln Lys Arg Asp Ala Asn Ser Ser Ile Tyr Lys Gly Lys 85 90 95

•

Lys Cys Arg M	et Glu 00	Ser Cys	Asp 105	Phe	Thr	Leu	Cys	Lys 110	Lys	Asn
-	00		100					110		

Gly	Phe	Lys	Val	Tyr	Val	Tyr	Pro	Gln	Gln	Lys	Gly	Glu	Lys	Ile	Ala
		115					120					125			

Glu Ser Tyr Gln Asn Ile Leu Ala Ala Ile Glu Gly Ser Arg Phe Tyr 130 135 140

Thr Ser Asp Pro Ser Gln Ala Cys Leu Phe Val Leu Ser Leu Asp Thr 145 150 150 160

Leu Asp Arg Asp Gln Leu Ser Pro Gln Tyr Val His Asn Leu Arg Ser 165 170 175

Lys Val Gln Ser Leu His Leu Trp Asn Asn Gly Arg Asn His Leu Ile 180 185 190

Phe Asn Leu Tyr Ser Gly Thr Trp Pro Asp Tyr Thr Glu Asp Val Gly 195 200 205

Phe Asp Ile Gly Gln Ala Met Leu Ala Lys Ala Ser Ile Ser Thr Glu 210 215 220

Asn Phe Arg Pro Asn Phe Asp Val Ser Ile Pro Leu Phe Ser Lys Asp 225 230 235 240

His Pro Arg Thr Gly Gly Glu Arg Gly Phe Leu Lys Phe Asn Thr Ile 245 250 255

Pro Pro Leu Arg Lys Tyr Met Leu Val Phe Lys Gly Lys Arg Tyr Leu 260 265 270

Thr Gly Ile Gly Ser Asp Thr Arg Asn Ala Leu Tyr His Val His Asn 275 280 285

Gly Glu	Asp	Val	Leu	Leu	Leu	Thr	Thr	Cys	Lys	His	Gly	Lys	Asp	\mathtt{Trp}
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Gln	Lys	His	Lys	Asp	Ser	Arg	Cys	Asp	Arg	Asp	Asn	Thr	Glu	Tyr	Glu
305	-		_		310					315					320

Leu	Leu	Leu	Gln	Ile	Pro	Ser	Thr	Ile	Arg	Ser	Ile	His	Gln	Asp	Lys
385					390					395					400

Thr Ala V	al Ile	His	Ala	Val	Thr	Pro	Leu	Val	Ser	Gln	Ser	Gln	Pro
		485					490			•		495	

Val	Leu	Lys	Leu	Leu	Val	Ala	Ala	Ala	Lys	Ser	Gln	Tyr	Cys	Ala	Gln
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Pro	Ala	Arg	Ser	His	Phe	Trp	Asp	Asn	Ser	Lys	Glu	Arg	Trp	Gly	Tyr
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Thr	Ser	Lys	Trp	Thr	Asn	Asp	Tyr	Ser	Met	Val	Leu	Thr	Gly	Ala	Ala
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Ala Ser Arg Trp Ala Asp Pro Asp His Phe Ala Gln Arg Gln Ser Cys 690 695 700

Met Asn Thr Phe Ala Ser Trp Phe Gly Tyr Met Pro Leu Ile His Ser 705 710 715 720

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<212> PRT

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Ser Ile Glu Ser Ser Ser Asp Gly Gly Val Glu Lys Arg Ser Ile Arg 50 55 60

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Gly	Phe	Asn	Pro 100	Lys	Asn	Lys	Ile	Lys 105	Val	Tyr	Ile	Tyr	Pro 110	Leu	Lys
Lys	Tyr	Val 115	Asp	Asp	Ala	Gly	Val 120	Pro	Val	Ser	Ser	Ala 125	Ile	Ser	Arg
Glu	Tyr 130	Asn	Glu	Leu	Leu	Thr 135	Ala	Ile	Ser	Asp	Ser 140	Asp	Tyr	Tyr	Thr
Asp 145	Asp	Ile	Asn	Arg	Ala 150	Cys	Leu	Phe	Val	Pro 155	Ser	Ile	Asp	Val	Leu 160
Asn	Gln	Asn	Pro	Leu 165	Arg	Ile	Lys	Glu	Thr 170	Ala	Gln	Ala	Leu	Ala 175	Gln
Leu	Ser	Arg	Trp 180	Asp	Arg	Gly	Thr	Asn 185	His	Leu	Leu	Phe	Asn 190	Met	Leu
Pro	Gly	Ala 195	Pro	Pro	Asp	Tyr	Asn 200	Thr	Ala	Leu	Asp	Val 205	Pro	Arg	Asp
Arg	Ala 210	Leu	Leu	Ala	Gly	Gly 215	Gly	Phe	Ser	Thr	Trp 220	Thr	Tyr	Arg	Gln
Gly 225	Tyr	Asp	Val	Ser	Ile 230	Pro	Val	Phe	Ser	Pro 235	Leu	Ser	Ala	Glu	Met 240
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250

255

Ala Leu Pro Glu Lys Ala Pro Gly Pro Arg Arg Tyr Phe Leu Leu Ser

245

Ser	Gln	Met	Ala	Ile	His	Pro	Glu	Tyr	Arg	Glu	Glu	Leu	Glu	Ala	Leu
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Leu Arg Arg Ala Arg Leu Gly Gln Ala Val Leu Ser Asp Val Leu Gln
$$325$$
 330 335

Lys	Met	Ser	Asp	Val	Tyr	Ser	Ile	Leu	Gln	Asn	Ile	Pro	Gln	Arg	Gln
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Ile	Pro	Pro	Gln	Ser	Gln	Gly	Phe	Thr	Ala	Ile	Val	Leu	Thr	Tyr	Asp
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Arg	Val	Glu	Ser	Leu	Phe	Arg	Val	Ile	Thr	Glu	Val	Ser	Lys	Val	Pro
465					470				•	475					480

Lys Ala Val Ile Lys Val Thr Pro Arg Lys Lys Phe Lys Cys Pro Glu 645 650 655

Cys Thr Ala Ile Asp Gly Leu Ser Leu Asp Gln Thr His Met Val Glu 660 665 670

Arg Ser Glu Cys Ile Asn Lys Phe Ala Ser Val Phe Gly Thr Met Pro 675 680 685

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Asp Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ile Ala Xaa Xaa Xaa Xaa Xaa 20 25 30

Val Xaa Xaa Xaa Xaa Xaa Xaa Asn Xaa Gly Xaa Tyr Xaa Xaa Xaa 35 40 45

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<223> all or part of sequence comprising residues 20-24 may be missing;
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<223> each position may be any amino acid

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each position may be any amino acid

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<213> Pasteurella multocida

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aataacttag aagttatcgt tgtagatgat tatagcacag ataaaacatt tcagatcgca 420

tccagaatag caaactctac aagtaaagta aaaacattcc gattaaactc aaatctaggg 480

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Page 163

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ctatgtaaga aaaacaatat actccaagtt tgtatatcac gaccatcgaa ttggctaaca Page 164

1680

gaagataaca aaaacactga gaccttattt catgaattcc aaaatagaga tgaaatacaa 1740

agtaaactca ttatttcaaa caacccttgg ggatactcaa gtatatatcc attattaaat 1800

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<210> 70

<211> 617

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<213> Pasteurella multocida

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Gln Ser Lys Ser Asn Lys Ile Glu Glu Asp Asn Ile Ser Gly Glu Asn 50 55 60

Lys Phe Ser Val Ser Ile Lys Asp Leu Tyr Asn Glu Ile Ser Asn Ser 65 70 75 80

Glu Leu Gly Ile Thr Lys Glu Arg Leu Gly Ala Pro Pro Leu Val Ser 85 90 95

Ile	Ile	Met	Thr	Ser	His	Asn	Thr	Glu	Lys	Phe	Ile	Glu	Ala	Ser	Ile
			100					105					110		

Asn	Ser	Leu	Leu	Leu	Gln	Thr	Tyr	Asn	Asn	Leu	Glu	Val	Ile	Val	Val
		115					120					125			

Cys	Ala	Tyr	Ser	Arg	Ile	Asn	Leu	Glu	Thr	Gln	Asn	Ile	Ile	Lys	Val
_	210					215					220				

Asn	Asp	Asn	Lys	Tyr	Lys	Leu	Gly	Leu	Ile	Thr	Leu	Gly	Val	Tyr	Arg
225					230					235					240

Ser	Leu	Phe	Ser	Asp	Met	Val	Glu	Trp	Val	Asp	Glu	Asn	Asn	Ile	Lys
	290					295					300				

Gln	Lys	Thr	Ser	Asp	Ala	Arg	Gln	Asn	Tyr	Leu	His	Glu	Phe	Gln	Lys
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